

Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1-14. (Cancelled)

15. (Currently amended) A method for synthesizing a double stranded nucleic acid molecule that contains information that represents computer-readable binary alphanumeric code, comprising at least the steps of:

- i) hybridizing together a plurality of double-stranded nucleic acid fragments, each fragment comprising at least one sequence of bases that represent a unit of the binary alphanumeric code, and each fragment comprising at least one single stranded region that is capable of hybridizing to at least one other fragment; and
- ii) optionally ligating the hybridized fragments; to produce a double stranded nucleic acid molecule comprising a series of binary alphanumeric code units.

16. (Previously presented) A method according to claim 15, wherein each sequence of bases that represents a unit of the alphanumeric code consists of between 4 and 10 bases.

17. (Previously presented) A method according to claim 15 or claim 16, wherein each fragment consists of between 8 and 25 bases.

18. (Cancelled)

19. (Previously presented) A method according to claim 15, wherein at least 10 double stranded nucleic acid fragments are hybridized together in step (i), to produce a double-stranded nucleic acid molecule comprising 10 fragments.

20. (Previously presented) A method according to claim 15, wherein a plurality of double stranded nucleic acid molecules comprising a series of double-stranded nucleic acid fragments are synthesized and linked together.

21. (Currently amended) A double stranded nucleic acid molecule that contains information that represents computer-readable binary alphanumeric code, produced according to the method according to claim 15.

22. (Currently amended) A method of identifying at least one binary alphanumeric code unit contained within a double stranded nucleic acid molecule produced according to claim 15, comprising the steps of:

- i) binding a labelled probe that is specific to at least one alphanumeric code unit to the unit; and

- ii) detecting the label associated with the bound probe,
thereby detecting the presence of the binary
alphanumeric code unit to which the probe binds.

23. (Previously presented) A library comprising a plurality of double stranded nucleic acid fragments as defined in claim 15.

24. (Currently amended) A kit for synthesizing a double stranded nucleic acid molecule that contains information that represents computer-readable binary alphanumeric code comprising a library of fragments according to claim 15 and a ligase.